

A DOCPHOENIX

APPL PARTS

IMIS
Internal Misc. Paper
LET.
Misc. Incoming Letter
371P
PCT Papers in a 371Application
Α
Amendment Including Elections
ABST
Abstract
ADS
Application Data Sheet
AF/D
Affidavit or Exhibit Received
APPENDIX
Appendix
ARTIFACT
Artifact
BIB
Bib Data Sheet
CLM
Claim
COMPUTER
Computer Program Listing
29.03.02 CRFL
All CRF Papers for Backfile
DIST
Terminal Disclaimer Filed
DRW
Drawings
FOR
Foreign Reference
FRPR
Foreign Priority Papers
IDS

NPL
Non-Patent Literature
OATH
Oath or Declaration
PET
Petition
RETMAIL
Mail Returned by USPS
SEQLIST
Sequence Listing
SPEC
Specification
SPEC NO
Specification Not in English
TRNA
Transmittal New Application

	CTNF
Count Non-Fina	al
On the Dankisti	CTRS
Count Restriction	on
	EXIN
Examiner Inter	view
	M903
DO/EO Accepta	ance
	M905
DO/EO Missing	Requirement
	NFDR
Formal Drawing	
	NOA
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	PETDEC
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OUTGOING

	CTMS
Misc. Office Ac	tion
	1449
Signed 1449	
	892
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	APEA
Examiner Answ	
	CTAV
Count Advisory	Action
	CTEQ
Count Ex parte	
	CTFR

Count Final Rejection

Appeal Brief
C.AD
Change of Address
N/AP
Notice of Appeal
PA
Change in Power of Attorney
REMApplicant Remarks in Amendment
XT/
Extension of Time filed concrete

INCOMING

File Wrapper

6/26/03

IDS Including 1449



PCT09

Does Not Comply Corrected Diskette Needed

Error on p. Z

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/936,845A

DATE: 03/29/2002 TIME: 13:38:02

Input Set : A:\09-936,845 substitute sequence listing.txt

Output Set: N:\CRF3\03292002\1936845A.raw

- 3 <110> APPLICANT: Napier, Johnathan A.
- 5 < 120 > TITLE OF INVENTION: Polyunsaturated Fatty Acid (PUFA) Elongase from Caenorhabditis elegans
 - 7 <130> FILE REFERENCE: 76/7
- C--> 9 <140> CURRENT APPLICATION NUMBER: US/09/936,845A
- C--> 10 <141> CURRENT FILING DATE: 2002-03-06
 - 12 <160> NUMBER OF SEQ ID NOS: 22
 - 14 <170> SOFTWARE: PatentIn Ver. 2.1

ERRORED SEQUENCES

- 728 <210> SEQ ID NO: 22
- 729 <211> LENGTH: 269
- 730 <212> TYPE: PRT
- 731 <213> ORGANISM: C. elegans
- 733 <220> FEATURE:
- 734 <221> NAME/KEY: VARIANT
- 735 <222> LOCATION: (1)...(269)
- 736 <223> OTHER INFORMATION: Xaa = Any Amino Acid
- 738 <400> SEQUENCE: 22
- 739 Met Ser Ala Glu Val Ser Glu Arg Phe Lys Val Trp Thr Gly Asn Asn 740
- 740 1 5 10 15
- 742 Glu Thr Ile Ile Tyr Ser Pro Phe Glu Tyr Asp Ser Thr Leu Leu Ile 743 20 25 30
- 745 Glu Ser Cys Arg Cys Thr Tyr Gln Leu Leu Ile Leu Leu Arg Gln Ile
- 746 35 40 45
 W--> 748 Tyr Tyr Arg Asp Ile Trp Ser His Gly Asn Leu Lys Ala Cys Asp Xaa
- 749 50 55 60
 - 751 Leu Leu Leu Ala Trp Asn Gly Phe Leu Ala Val Phe Ser Ile Met Gly 752 65 70 75 80
- W--> 754 Thr Trp Arg Phe Gly Ile Glu Phe Tyr Asp Ala Val Phe Arg Xaa Gly
 755 85 90 95
- W--> 757 Phe Ile Xaa Ser Ile Cys Leu Ala Val Asn Pro Arg Ser Pro Ser Ala
 758 100 105 110
 - 760 Phe Trp Ala Cys Met Phe Ala Leu Ser Lys Ile Ala Glu Phe Gly Asp 761 115 120 125
 - 763 Thr Met Phe Leu Val Leu Arg Lys Arg Pro Val Ile Phe Leu His Trp
 - 764 130 135 140 766 Tyr His His Ala Val Val Leu Ile Leu Ser Trp His Ala Ala Ile Glu
 - 767 145 150 150 155 160
 - 769 Leu Thr Ala Pro Gly Arg Trp Phe Ile Phe Met Asn Tyr Leu Val His 770 165 170 175
- W--> 772 Ser Ile Met Tyr Thr Tyr Tyr Ala Ile Thr Ser Ile Gly Tyr Arg Xaa

RAW SEQUENCE LISTING DATE: 03/29/2002 PATENT APPLICATION: US/09/936,845A TIME: 13:38:03

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Output Set: N:\CRF3\03292002\1936845A.raw

185 180 775 Pro Lys Ile Val Ser Met Thr Val Thr Phe Leu Gln Thr Leu Gln Met 195 776 778 Leu Ile Gly Val Ser Ile Ser Cys Ile Val Leu Tyr Leu Lys Leu Asn 215 220 781 Gly Glu Met Cys Gln Gln Ser Tyr Asp Asn Leu Ala Leu Ser Phe Gly 235 230 784 Ile Tyr Ala Ser Phe Leu Val Leu Ser Ser Phe Phe Asn Asn Ala Tyr 250 245 787 Leu Val Lys Lys Asp Lys Lys Pro Asp Val Lys Lys Asp 788 260 E--> 792(1